

GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: August 12, 2004, 15:34:29 ; Search time 0.001 Seconds  
(without alignments)  
164.450 Million cell updates/sec

Title: US-10-033-742-3

Perfect score: 65

Sequence: 1 ttttgaatggaattggac.....gctggggttgagggtttac 65

Scoring table: IDENTITY\_NUC

Gapop 10.0 , Gapext 0.5

Searched: 103 seqs, 1265 residues

Total number of hits satisfying chosen parameters: 206

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 103 summaries

Database : rndb.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
C 1	19.8	30.5	23	1	US-09-071-353-12
C 2	19.8	30.5	23	1	US-09-426-326-12
C 3	14	21.5	18	1	US-08-525-654A-138
C 4	12.8	19.7	17	1	US-08-281-940-29
C 5	12.8	19.7	17	1	US-08-710-134-29
C 6	12.8	19.7	17	1	US-08-485-885-29
C 7	12.8	19.7	17	1	US-09-866-108A-2464
C 8	12.8	19.7	17	1	US-09-866-108A-2465
C 9	11.4	17.5	15	1	US-08-146-886-12
C 10	11.4	17.5	15	1	US-08-440-787A-139
C 11	11.4	17.5	15	1	US-09-109-613-22
C 12	11.4	17.5	15	1	US-08-730-635-5
C 13	11.4	17.5	15	1	US-08-730-635-9
C 14	10.8	16.6	14	1	US-08-242-664-25
C 15	10.8	16.6	14	1	US-08-484-138-25
C 16	10.8	16.6	14	1	US-09-580-923-29
C 17	10.8	16.6	14	1	US-09-580-923-30
C 18	10.8	16.6	14	1	PCT-US95-06379-25
C 19	10.4	16.0	12	1	US-08-004-800-9
C 20	10.4	16.0	12	1	US-08-004-800-11
C 21	10.4	16.0	12	1	US-08-004-800-12
C 22	10.4	16.0	12	1	US-08-115-497-14
C 23	10.4	16.0	12	1	US-08-115-497-15
C 24	10.4	16.0	12	1	US-08-115-497-17
C 25	10.4	16.0	12	1	US-08-413-813-9
C 26	10.4	16.0	12	1	US-08-413-813-10
C 27	10.4	16.0	12	1	US-08-413-813-11
C 28	10.4	16.0	12	1	US-08-413-813-12
C 29	10.4	16.0	12	1	US-08-413-813-13
C 30	10.4	16.0	12	1	US-08-413-813-14
C 31	10.4	16.0	12	1	US-08-466-670-14
C 32	10.4	16.0	12	1	US-08-466-670-15
C 33	10.4	16.0	12	1	US-08-466-670-17

## ALIGNMENTS

34	10.4	16.0	12	1	US-08-467-346-9	Sequence 9, Appl
C 35	10.4	16.0	12	1	US-08-467-346-10	Sequence 10, Appl
C 36	10.4	16.0	12	1	US-08-467-346-28	Sequence 28, Appl
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C 38	10.4	16.0	12	1	US-08-467-346-31	Sequence 31, Appl
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C 40	10.4	16.0	12	1	PCT-US92-02480A-9	Sequence 9, Appl
C 41	10.4	16.0	12	1	PCT-US92-02480A-11	Sequence 11, Appl
C 42	10.4	16.0	12	1	PCT-US92-02480A-12	Sequence 12, Appl
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C 57	9.4	14.5	12	1	US-08-466-670-13	Sequence 13, Appl
C 58	9.4	14.5	12	1	US-08-494-301A-12	Sequence 12, Appl
C 59	9.4	14.5	12	1	US-08-467-346-38	Sequence 38, Appl
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C 61	9.4	14.5	12	1	US-08-403-888A-41	Sequence 41, Appl
C 62	9.4	14.5	12	1	US-08-403-888A-57	Sequence 57, Appl
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C 68	9.4	14.5	12	1	US-09-378-535-5	Sequence 5, Appl
C 69	9.4	14.5	12	1	US-09-378-535-33	Sequence 33, Appl
C 70	9.4	14.5	12	1	US-09-378-535-35	Sequence 35, Appl
C 71	9.4	14.5	12	1	PCT-US94-02471-53	Sequence 53, Appl
C 72	9.4	14.5	12	1	US-08-482-115B-32	Sequence 32, Appl
C 73	9	13.8	9	1	US-08-472-802C-32	Sequence 32, Appl
C 74	9	13.8	9	1	US-09-057-351-32	Sequence 32, Appl
C 75	9	13.8	9	1	US-08-330-123A-10	Sequence 10, Appl
C 76	9	13.8	10	1	US-08-482-115B-10	Sequence 10, Appl
C 77	9	13.8	10	1	US-08-660-678A-10	Sequence 10, Appl
C 78	9	13.8	10	1	US-08-485-708-41	Sequence 41, Appl
C 79	9	13.8	10	1	US-08-472-802C-11	Sequence 11, Appl
C 80	9	13.8	10	1	US-08-388-353-513	Sequence 513, Appl
C 81	9	13.8	10	1	US-08-388-353-514	Sequence 514, Appl
C 82	9	13.8	10	1	US-08-388-353-547	Sequence 547, Appl
C 83	9	13.8	10	1	US-08-488-551B-513	Sequence 513, Appl
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C 98	9	13.8	12	1	PCT-US96-09430-17	Sequence 17, Appl
C 99	9	13.8	12	1	PCT-US96-09430-18	Sequence 18, Appl

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OM nucleic - nucleic search, using sw model

Run on: August 12, 2004, 15:30:40 ; Search time 1 Seconds  
(without alignments)  
0.117 Million cell updates/sec

Title: US-10-033-742-3  
Perfect score: 65  
Sequence: 1 tttcgggaatggaatggac.....gcctggggtggaggttccac 65

Scoring table: IDENTITY NUC  
Gapop 10.0 , Gapext 0.5

Searched: 61 seqs, 898 residues

Total number of hits satisfying chosen parameters: 122

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 61 summaries

Database : pub:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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6	20	30.8	20	1	US-10-033-742-23
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18	12.8	19.7	17	1	US-09-866-108-2465
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*Published - Applications - NA*

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C 38	10.4	16.0	13	1	US-10-253-904-45	Sequence 45, Appl
C 39	10.4	16.0	13	1	US-10-392-970-36	Sequence 36, Appl
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C 41	10	15.4	10	1	US-10-330-627-973	Sequence 973, App
C 42	10	15.4	10	1	US-10-450-797-847	Sequence 847, App
C 43	10	15.4	12	1	US-10-091-281-67	Sequence 67, Appl
C 44	9.8	15.1	13	1	US-09-789-836-31	Sequence 31, Appl
C 45	9.8	15.1	13	1	US-09-789-836-31	Sequence 31, Appl
C 46	9.4	14.5	11	1	US-09-918-715-8	Sequence 8, Appl
C 47	9.4	14.5	11	1	US-10-450-797-174	Sequence 174, App
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C 61	9.4	14.5	12	1	US-10-422-262-22	Sequence 22, Appl

## ALIGNMENTS

## RESULT 1

US-10-717-597-3888  
; Sequence 3888, Application US/10717597  
; Publication No. US20040110221A1  
; GENERAL INFORMATION:  
; APPLICANT: Wyeth  
; APPLICANT: Burczynski, Michael E.  
; APPLICANT: Twine, Natalie C.  
; APPLICANT: Dorne, Andrew J.  
; APPLICANT: Trepicchio, William L.  
; APPLICANT: Slonim, Donna K.  
; APPLICANT: Stover, Jennifer A.  
; TITLE OF INVENTION: METHODS FOR DIAGNOSING RCC AND OTHER SOLID TUMORS  
; FILE REFERENCE: AM101080L  
; CURRENT APPLICATION NUMBER: US/10717,597  
; PRIOR FILING DATE: 2003-11-21  
; PRIOR FILING DATE: 2003-11-21  
; PRIOR FILING DATE: 2003-04-03  
; PRIOR FILING DATE: 2003-04-03  
; PRIOR FILING DATE: 2002-11-21  
; NUMBER OF SEQ ID NOS: 4904  
; SOFTWARE: Patent in version 3.2  
; SEQ ID NO 3888  
; LENGTH: 25  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-10-717-597-3888

Query Match 38.5%; Score 25; DB 1; Length 25;  
Best Local Similarity 100.0%; Pred. No. 0.62;  
Matches 25; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 1 GGAATTGGACATAGCCCAAGACAG 25

## RESULT 2

US-10-717-597-3889  
; Sequence 3889, Application US/10717597

## ATTACHMENTS

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/mol_type="unassigned DNA"
db_xref="taxon:32644"

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DEFINITION	AR300570					
ACCESSION	AR300570.1	GI:31688075				
VERSION						
KEYWORDS						
SOURCE	Unknown.					
ORGANISM	Unknown.					
	Unclassified.					

Query Match	30.5%;	Score 19.8;	DB 1;	Length 23;
Best Local Similarity	91.3%;	Pred. No. 1.6;		
Mismatches	0;	Mismatches	2;	Indels
Conservative	0;	Gaps	0;	Gaps

RESULT 4	21 bp	DNA	linear	PAT 28-FEB-2001
AX084313/c	AX084313			
LOCUS	LOCUS			
SEQUENCE	Sequence	107	from Patent	WO0110902.

ACCESSION  
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 VERSION  
 AX084333.1 GI:13185815  
 KEYWORDS  
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 SOURCE  
 synthetic construct  
 ORGANISM  
 artificial sequences.  
 1  
 REFERENCE  
 1  
 AUTHORS  
 Shinkets, R. A. and Fernandes, E.  
 TITLE  
 Nucleic acids and secreted polypeptides encoded thereby

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OM nucleic - nucleic search, using sw model

Run on: August 12, 2004, 15:32:42 ; Search time 1 Seconds  
(without alignments)  
0.227 Million cell updates/sec

Title: US-10-033-742-3

Perfect score: 65

Sequence: 1 ttctggaatggattggac.....gctgggggttggaaggtttcac 65

Scoring table: IDENTITY\_NUC

Gapop 10.0 , Gapext 0.5

Searched: 134 seqs, 1745 residues

Total number of hits satisfying chosen parameters: 268

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 134 summaries

Database : rge.db:\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

# SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
C 1	19.8	30.5	23	1	ACCESSION: A83656
C 2	19.8	30.5	23	1	ACCESSION: AR300570
C 3	19.8	30.5	23	1	ACCESSION: BD106469
C 4	15.8	24.3	21	1	ACCESSION: AX084313
C 5	14.4	22.2	18	1	ACCESSION: BD107600
C 6	14	21.5	18	1	ACCESSION: AR000413
C 7	13.8	21.2	17	1	ACCESSION: BD183671
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C 9	13.4	20.6	17	1	ACCESSION: AX217387
C 10	13	20.0	17	1	ACCESSION: AX217388
C 11	13	20.0	17	1	ACCESSION: AX217388
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C 13	13	20.0	17	1	ACCESSION: AX733882
C 14	12.8	19.7	17	1	ACCESSION: AX757493
C 15	12.8	19.7	17	1	ACCESSION: AR053059
C 16	12.8	19.7	17	1	ACCESSION: AR055020
C 17	12.8	19.7	17	1	ACCESSION: BD254337
C 18	12.8	19.7	17	1	ACCESSION: BD258370
C 19	12.8	19.7	17	1	ACCESSION: BD258370
C 20	12.4	19.1	15	1	ACCESSION: AX218301
C 21	12.4	19.1	15	1	ACCESSION: BD233057
C 22	12.4	19.1	15	1	ACCESSION: BD233078
C 23	12.4	19.1	15	1	ACCESSION: AX007611
C 24	11.4	17.5	13	1	ACCESSION: AX007632
C 25	11.4	17.5	13	1	ACCESSION: AX104604
C 26	11.4	17.5	13	1	ACCESSION: AX355422
C 27	11.4	17.5	14	1	ACCESSION: AX547657
C 28	11.4	17.5	14	1	ACCESSION: BD233079
C 29	11.4	17.5	15	1	ACCESSION: AX007633
C 30	11.4	17.5	15	1	ACCESSION: I45950
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39	10.8	16.6	14	1	I28572
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42	10.8	16.6	14	1	AX323394
C 43	10.8	16.6	14	1	AX323395
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C 84	10	15.4	12	1	AR349261
C 85	9.8	15.1	13	1	A01985
C 86	9.8	15.1	13	1	A06431
C 87	9.8	15.1	13	1	BD062265
88	9.4	14.5	11	1	I16094
C 89	9.4	14.5	11	1	AX393078
C 90	9.4	14.5	11	1	AX470597
C 91	9.4	14.5	11	1	AX470878
C 92	9.4	14.5	11	1	AX471365
C 93	9.4	14.5	11	1	AX623489
C 94	9.4	14.5	11	1	AX624179
C 95	9.4	14.5	11	1	AX624329
96	9.4	14.5	11	1	AX624484
C 97	9.4	14.5	11	1	AX625720
C 98	9.4	14.5	11	1	AX625789
C 99	9.4	14.5	11	1	AX626474
C 100	9.4	14.5	11	1	AX627570
C 101	9.4	14.5	11	1	AX628639
C 102	9.4	14.5	11	1	AX628640
C 103	9.4	14.5	11	1	AX628771
C 104	9.4	14.5	11	1	AX629905
C 105	9.4	14.5	11	1	AX630278
C 106	9.4	14.5	11	1	AX630910

GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: August 12, 2004, 15:28:53 ; Search time 1.2 seconds  
(without alignments)  
0.904 Million cell updates/sec

Title: US-10-033-742-3  
Perfect score: 65  
Sequence: 1 ttcttggaatggaattggac.....gctggggttgagggtttcac 65

Scoring table: IDENTITY\_NUC

Gapop 10.0 , Gapext 0.5

Searched: 534 seqs, 6954 residues

Total number of hits satisfying chosen parameters: 1068

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 534 summaries

Database : ngs:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

Result No.	Score	Query Match %	Length	ID	Description
C 1	24	36.9	24	1	ADC02407 Human macrophage i
C 2	20	30.8	20	1	AD57273 Human MIP3A DNA sp
C 3	20	30.8	20	1	AD57273 Human MIP3A DNA sp
C 4	20	30.8	20	1	AD57276 Human MIP3A DNA sp
C 5	20	30.8	20	1	AD57275 Human MIP3A DNA sp
C 6	19.8	30.5	23	1	AAV82678 Biotinylated probe
C 7	15.8	24.3	21	1	AAF74520 Clone 16467945 PRO
C 8	14.4	22.2	18	1	AA42340 Novel sand pear mi
C 9	14	21.5	18	1	AAQ99361 Japanese oyster tr
C 10	13.4	20.6	16	1	ABX75321 Human 216 Gene all
C 11	13.4	20.6	17	1	ABX02829 Human CD20 Hammer
C 12	13.4	20.6	17	1	ABX02828 Human CD20 Hammer
C 13	13	20.0	17	1	ABX03198 Human CD20 Hammer
C 14	13	20.0	17	1	ABX03198 Human CD20 Hammer
C 15	13	20.0	17	1	ABT39879 Human CD20 Inozyme
C 16	13	20.0	17	1	AD840491 Tumour suppression
C 17	12.8	19.7	17	1	AD60238 Tumour suppression
C 18	12.8	19.7	17	1	AAQ93361 ASO Q493XM repress
C 19	12.8	19.7	17	1	AAQ93361 ASO Q493XM repress
C 20	12.8	19.7	17	1	AAQ93361 ASO Q493XM repress
C 21	12.8	19.7	17	1	AAQ93361 ASO Q493XM repress
C 22	12.8	19.7	17	1	AAQ93361 ASO Q493XM repress
C 23	12.4	19.1	15	1	AAQ93361 ASO Q493XM repress
C 24	12.4	19.1	15	1	AAQ93361 ASO Q493XM repress
C 25	12	18.5	12	1	AB167380 HIV-1 protease gen
C 26	12	18.5	12	1	AB167380 HIV-1 protease gen
C 27	12	18.5	12	1	AB167380 HIV-1 protease gen
C 28	12	18.5	12	1	AB167380 HIV-1 protease gen
C 29	12	18.5	12	1	AB167380 HIV-1 protease gen
C 30	11.8	18.2	15	1	AAQ93361 ASO Q493XM repress
C 31	11.8	18.2	15	1	AAQ93361 ASO Q493XM repress
C 32	11.8	18.2	15	1	AAQ93361 ASO Q493XM repress
C 33	11.8	18.2	15	1	AAQ93361 ASO Q493XM repress

34	11.4	17.5	13	1	AAF99594 Immunostimulatory
35	11.4	17.5	13	1	ABF69510 Oligonucleotide SE
C 36	11.4	17.5	13	1	ABH44501 Oligonucleotide SE
C 37	11.4	17.5	13	1	ABF69511 Oligonucleotide SE
C 38	11.4	17.5	13	1	ABF07727 Oligonucleotide SE
C 39	11.4	17.5	13	1	ABC73222 Oligonucleotide SE
C 40	11.4	17.5	13	1	ABC73223 Oligonucleotide SE
41	11.4	17.5	13	1	ABF07726 Oligonucleotide SE
42	11.4	17.5	13	1	ABH57826 Oligonucleotide SE
43	11.4	17.5	13	1	ABH44500 Oligonucleotide SE
C 44	11.4	17.5	13	1	ABH57827 Oligonucleotide SE
45	11.4	17.5	13	1	ABF78312 Angiogenesis inh
46	11.4	17.5	13	1	ABL39046 Immunostimulatory
47	11.4	17.5	13	1	ACH03134 Immunostimulatory
48	11.4	17.5	13	1	ADB37036 Immunostimulatory
49	11.4	17.5	14	1	AAZ97685 HIV-1 protease Gen
50	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 51	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
52	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
53	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 54	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 55	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
56	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 57	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
58	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 59	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
60	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
61	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
62	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
63	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
64	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 65	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
66	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
67	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 68	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 69	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
70	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 71	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 72	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
73	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
C 74	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
75	11.4	17.5	15	1	AAQ90131 69-mer oligonucleo
76	10.8	16.6	14	1	AAQ90131 69-mer oligonucleo
77	10.8	16.6	14	1	AAQ90131 69-mer oligonucleo
78	10.8	16.6	14	1	AAQ90131 69-mer oligonucleo
C 79	10.8	16.6	14	1	AAQ90131 69-mer oligonucleo
C 80	10.6	16.3	13	1	AAQ90131 69-mer oligonucleo
81	10.6	16.3	13	1	AAQ90131 69-mer oligonucleo
C 82	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 83	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
84	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
85	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 86	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
87	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 88	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 89	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 90	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
91	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 92	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
93	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 94	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
95	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
96	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 97	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
98	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 99	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
100	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 101	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
102	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 103	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
104	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
C 105	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo
106	10.4	16.0	12	1	AAQ90131 69-mer oligonucleo

*N. G. Gendley*

c 107	107	10.4	16.0	12	1	AB124669	Oligonucleotide pr
c 108	108	10.4	16.0	12	1	AB157127	Oligonucleotide pr
c 109	109	10.4	16.0	12	1	ABH70112	Oligonucleotide pr
c 110	110	10.4	16.0	12	1	AB175408	Oligonucleotide pr
c 111	111	10.4	16.0	12	1	ABH75819	Oligonucleotide pr
c 112	112	10.4	16.0	12	1	AB108168	Oligonucleotide pr
c 113	113	10.4	16.0	12	1	AB111964	Oligonucleotide pr
c 114	114	10.4	16.0	12	1	AB111865	Oligonucleotide pr
c 115	115	10.4	16.0	12	1	AB113865	Oligonucleotide pr
c 116	116	10.4	16.0	12	1	AB138926	Oligonucleotide pr
c 117	117	10.4	16.0	12	1	ABH73791	Oligonucleotide pr
c 118	118	10.4	16.0	12	1	AB123551	Oligonucleotide pr
c 119	119	10.4	16.0	12	1	AB142492	Oligonucleotide pr
c 120	120	10.4	16.0	12	1	AB175632	Oligonucleotide pr
c 121	121	10.4	16.0	12	1	AB128119	Oligonucleotide pr
c 122	122	10.4	16.0	12	1	AB117994	Oligonucleotide pr
c 123	123	10.4	16.0	12	1	AB107453	Oligonucleotide pr
c 124	124	10.4	16.0	12	1	AB162010	Oligonucleotide pr
c 125	125	10.4	16.0	12	1	AB100155	Oligonucleotide pr
c 126	126	10.4	16.0	13	1	ABC95493	Oligonucleotide SE
c 127	127	10.4	16.0	13	1	ABC72046	Oligonucleotide SE
c 128	128	10.4	16.0	13	1	ABC11945	Oligonucleotide SE
c 129	129	10.4	16.0	13	1	ABC87797	Oligonucleotide SE
c 130	130	10.4	16.0	13	1	ABC89082	Oligonucleotide SE
c 131	131	10.4	16.0	13	1	ABC60882	Oligonucleotide SE
c 132	132	10.4	16.0	13	1	ABF09494	Oligonucleotide SE
c 133	133	10.4	16.0	13	1	ABH08614	Oligonucleotide SE
c 134	134	10.4	16.0	13	1	ABC18620	Oligonucleotide SE
c 135	135	10.4	16.0	13	1	ABC11997	Oligonucleotide SE
c 136	136	10.4	16.0	13	1	ABC92634	Oligonucleotide SE
c 137	137	10.4	16.0	13	1	ABC22672	Oligonucleotide SE
c 138	138	10.4	16.0	13	1	ABF53874	Oligonucleotide SE
c 139	139	10.4	16.0	13	1	ABC93477	Oligonucleotide SE
c 140	140	10.4	16.0	13	1	ABC19996	Oligonucleotide SE
c 141	141	10.4	16.0	13	1	ABP09495	Oligonucleotide SE
c 142	142	10.4	16.0	13	1	ABH45216	Oligonucleotide SE
c 143	143	10.4	16.0	13	1	ABF12720	Oligonucleotide SE
c 144	144	10.4	16.0	13	1	ABH18990	Oligonucleotide SE
c 145	145	10.4	16.0	13	1	ABF72258	Oligonucleotide SE
c 146	146	10.4	16.0	13	1	ABH64846	Oligonucleotide SE
c 147	147	10.4	16.0	13	1	ABC61688	Oligonucleotide SE
c 148	148	10.4	16.0	13	1	ABH18991	Oligonucleotide SE
c 149	149	10.4	16.0	13	1	ABC95492	Oligonucleotide SE
c 150	150	10.4	16.0	13	1	ABC87700	Oligonucleotide SE
c 151	151	10.4	16.0	13	1	ABF32872	Oligonucleotide SE
c 152	152	10.4	16.0	13	1	ABH56067	Oligonucleotide SE
c 153	153	10.4	16.0	13	1	ABC69962	Oligonucleotide SE
c 154	154	10.4	16.0	13	1	ABC00654	Oligonucleotide SE
c 155	155	10.4	16.0	13	1	ABP44599	Oligonucleotide SE
c 156	156	10.4	16.0	13	1	ABF72259	Oligonucleotide SE
c 157	157	10.4	16.0	13	1	ABC9476	Oligonucleotide SE
c 158	158	10.4	16.0	13	1	ABC00655	Oligonucleotide SE
c 159	159	10.4	16.0	13	1	ABF15052	Oligonucleotide SE
c 160	160	10.4	16.0	13	1	ABF15053	Oligonucleotide SE
c 161	161	10.4	16.0	13	1	ABF53875	Oligonucleotide SE
c 162	162	10.4	16.0	13	1	ABH08607	Oligonucleotide SE
c 163	163	10.4	16.0	13	1	ABH56066	Oligonucleotide SE
c 164	164	10.4	16.0	13	1	ABC9883	Oligonucleotide SE
c 165	165	10.4	16.0	13	1	ABC97701	Oligonucleotide SE
c 166	166	10.4	16.0	13	1	ABF67787	Oligonucleotide SE
c 167	167	10.4	16.0	13	1	ABF51528	Oligonucleotide SE
c 168	168	10.4	16.0	13	1	ABC69963	Oligonucleotide SE
c 169	169	10.4	16.0	13	1	ABC22673	Oligonucleotide SE
c 170	170	10.4	16.0	13	1	ABC95008	Oligonucleotide SE
c 171	171	10.4	16.0	13	1	ABC87800	Oligonucleotide SE
c 172	172	10.4	16.0	13	1	ABC98083	Oligonucleotide SE
c 173	173	10.4	16.0	13	1	ABF67787	Oligonucleotide SE
c 174	174	10.4	16.0	13	1	ABF51528	Oligonucleotide SE
c 175	175	10.4	16.0	13	1	ABC69963	Oligonucleotide SE
c 176	176	10.4	16.0	13	1	ABC22673	Oligonucleotide SE
c 177	177	10.4	16.0	13	1	ABC95008	Oligonucleotide SE
c 178	178	10.4	16.0	13	1	ABC87800	Oligonucleotide SE
c 179	179	10.4	16.0	13	1	ABC98083	Oligonucleotide SE
c 180	180	10.4	16.0	13	1	ABH08615	Oligonucleotide SE
c 181	181	10.4	16.0	13	1	AB18621	Oligonucleotide SE
c 182	182	10.4	16.0	13	1	ABF12721	Oligonucleotide SE
c 183	183	10.4	16.0	13	1	ABC87795	Oligonucleotide SE
c 184	184	10.4	16.0	13	1	ABF52773	Oligonucleotide SE
c 185	185	10.4	16.0	13	1	ABF44598	Oligonucleotide SE
c 186	186	10.4	16.0	13	1	ABC75858	Oligonucleotide SE
c 187	187	10.4	16.0	13	1	ABC1689	Oligonucleotide SE
c 188	188	10.4	16.0	13	1	ABF51529	Oligonucleotide SE
c 189	189	10.4	16.0	13	1	ABH08606	Oligonucleotide SE
c 190	190	10.4	16.0	13	1	ABH45217	Oligonucleotide SE
c 191	191	10.4	16.0	13	1	ABH4847	Oligonucleotide SE
c 192	192	10	15.4	10	1	AAQ79357	Sequence of lympho
c 193	193	10	15.4	10	1	AAV50176	Yeast tag for addi
c 194	194	10	15.4	10	1	AAZ77894	Human dendritic ce
c 195	195	10	15.4	10	1	AAZ82445	Metastatic breast
c 196	196	10	15.4	10	1	AAZ56570	Human macrophage
c 197	197	10	15.4	10	1	AAH41133	Human ubiquitously
c 198	198	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 199	199	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 200	200	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 201	201	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 202	202	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 203	203	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 204	204	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 205	205	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 206	206	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 207	207	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 208	208	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 209	209	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 210	210	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 211	211	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 212	212	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 213	213	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 214	214	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 215	215	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 216	216	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 217	217	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 218	218	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 219	219	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 220	220	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 221	221	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 222	222	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 223	223	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 224	224	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 225	225	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 226	226	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 227	227	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 228	228	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 229	229	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 230	230	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 231	231	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 232	232	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 233	233	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 234	234	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 235	235	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 236	236	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 237	237	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 238	238	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 239	239	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 240	240	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 241	241	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 242	242	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 243	243	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 244	244	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 245	245	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 246	246	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 247	247	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 248	248	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 249	249	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 250	250	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 251	251	10	15.4	10	1	AAH41485	Yeast NORF gene SA
c 252	252	10	15.4	10	1	AAH41485	Yeast NORF gene SA

253	10	15.4	13	1	ABP78290	Oligonucleotide SE
254	10	15.4	13	1	ABH09786	Oligonucleotide SE
C 255	10	15.4	13	1	ABH14635	Oligonucleotide SE
C 256	10	15.4	13	1	ABF35131	Oligonucleotide SE
257	10	15.4	13	1	ABF68180	Oligonucleotide SE
258	10	15.4	13	1	ABF72180	Oligonucleotide SE
C 259	10	15.4	13	1	ABF72191	Oligonucleotide SE
260	10	15.4	13	1	ABF78294	Oligonucleotide SE
261	10	15.4	13	1	ABH12398	Oligonucleotide SE
262	10	15.4	13	1	ABH12390	Oligonucleotide SE
C 263	10	15.4	13	1	ABC18903	Oligonucleotide SE
C 264	10	15.4	13	1	ABC69993	Oligonucleotide SE
C 265	10	15.4	13	1	ABC53076	Oligonucleotide SE
266	10	15.4	13	1	ABF14904	Oligonucleotide SE
C 267	10	15.4	13	1	ABC16841	Oligonucleotide SE
268	10	15.4	13	1	ABF29456	Oligonucleotide SE
C 269	10	15.4	13	1	ABF78295	Oligonucleotide SE
C 270	10	15.4	13	1	ABH28568	Oligonucleotide SE
C 271	10	15.4	13	1	ABH09787	Oligonucleotide SE
C 272	10	15.4	13	1	ABH41477	Oligonucleotide SE
C 273	10	15.4	13	1	ABC38819	Oligonucleotide SE
274	10	15.4	13	1	ABF35110	Oligonucleotide SE
C 275	10	15.4	13	1	ABF68181	Oligonucleotide SE
C 276	10	15.4	13	1	ABF72191	Oligonucleotide SE
C 277	10	15.4	13	1	ABH28569	Oligonucleotide SE
278	10	15.4	13	1	ABH66664	Oligonucleotide SE
C 279	10	15.4	13	1	ABC08153	Oligonucleotide SE
C 280	10	15.4	13	1	ABC08156	Oligonucleotide SE
C 281	10	15.4	13	1	ABC8851	Oligonucleotide SE
C 282	10	15.4	13	1	ABF14836	Oligonucleotide SE
C 283	10	15.4	13	1	ABF39155	Oligonucleotide SE
284	10	15.4	13	1	ABH36930	Oligonucleotide SE
285	10	15.4	13	1	ABH59464	Oligonucleotide SE
286	10	15.4	13	1	ABF21342	Oligonucleotide SE
C 287	10	15.4	13	1	ABH27163	Oligonucleotide SE
C 288	10	15.4	13	1	ABF52228	Oligonucleotide SE
289	10	15.4	13	1	ABF60990	Oligonucleotide SE
C 290	10	15.4	13	1	ABC68992	Oligonucleotide SE
C 291	10	15.4	13	1	ABF02807	Oligonucleotide SE
292	10	15.4	13	1	ABC66872	Oligonucleotide SE
C 293	10	15.4	13	1	ABF18949	Oligonucleotide SE
C 294	10	15.4	13	1	ABF24945	Oligonucleotide SE
C 295	10	15.4	13	1	ABF24945	Oligonucleotide SE
C 296	10	15.4	13	1	ABF73195	Oligonucleotide SE
297	10	15.4	13	1	ABC69992	Oligonucleotide SE
C 298	10	15.4	13	1	ABF18948	Oligonucleotide SE
C 299	10	15.4	13	1	ABH16777	Oligonucleotide SE
C 300	10	15.4	13	1	ABH42921	Oligonucleotide SE
301	10	15.4	13	1	ABF14837	Oligonucleotide SE
C 302	10	15.4	13	1	ABP21338	Oligonucleotide SE
C 303	10	15.4	13	1	ABP21339	Oligonucleotide SE
C 304	10	15.4	13	1	ABF39359	Oligonucleotide SE
C 305	10	15.4	13	1	ABH36931	Oligonucleotide SE
C 306	10	15.4	13	1	ABH46665	Oligonucleotide SE
C 307	10	15.4	13	1	ABC08157	Oligonucleotide SE
C 308	10	15.4	13	1	ABP21343	Oligonucleotide SE
309	10	15.4	13	1	ABF24944	Oligonucleotide SE
C 310	10	15.4	13	1	ABF52229	Oligonucleotide SE
C 311	10	15.4	13	1	ABH10859	Oligonucleotide SE
C 312	10	15.4	13	1	ABH12999	Oligonucleotide SE
313	10	15.4	13	1	ABH41476	Oligonucleotide SE
314	10	15.4	13	1	ABF14180	Oligonucleotide SE
C 315	10	15.4	13	1	ABC66873	Oligonucleotide SE
316	10	15.4	13	1	ABF39154	Oligonucleotide SE
C 317	10	15.4	13	1	ABF39015	Oligonucleotide SE
C 318	10	15.4	13	1	ABF52057	Oligonucleotide SE
319	10	15.4	13	1	ABH14634	Oligonucleotide SE
C 320	10	15.4	13	1	ABH59465	Oligonucleotide SE
C 321	10	15.4	13	1	ABC68993	Oligonucleotide SE
C 322	10	15.4	13	1	ABF29457	Oligonucleotide SE
323	10	15.4	13	1	ABF29458	Oligonucleotide SE
C 324	10	15.4	13	1	ABC74005	Oligonucleotide SE
325	10	15.4	13	1	ABC74005	Oligonucleotide SE

[illegible]

C 399	9.8	15.1	13	1	ABF47357	Oligonucleotide SE
400	9.8	15.1	13	1	ABF33692	Oligonucleotide SE
401	9.8	15.1	13	1	ABH33542	Oligonucleotide SE
C 402	9.8	15.1	13	1	ABF58451	Oligonucleotide SE
403	9.8	15.1	13	1	ABH34278	Oligonucleotide SE
404	9.8	15.1	13	1	ABF86466	Oligonucleotide SE
C 405	9.8	15.1	13	1	ABH16219	Oligonucleotide SE
406	9.8	15.1	13	1	ABH62129	Oligonucleotide SE
C 407	9.8	15.1	13	1	ABC31789	Oligonucleotide SE
408	9.8	15.1	13	1	ABF33180	Oligonucleotide SE
C 409	9.8	15.1	13	1	ABF57881	Oligonucleotide SE
C 410	9.8	15.1	13	1	ABH04581	Oligonucleotide SE
C 411	9.8	15.1	13	1	ABH05560	Oligonucleotide SE
C 412	9.8	15.1	13	1	ABC74671	Oligonucleotide SE
C 413	9.8	15.1	13	1	ABF04647	Oligonucleotide SE
414	9.8	15.1	13	1	ABF07730	Oligonucleotide SE
C 415	9.8	15.1	13	1	ABC34457	Oligonucleotide SE
416	9.8	15.1	13	1	ABC60968	Oligonucleotide SE
C 417	9.8	15.1	13	1	ABF20477	Oligonucleotide SE
418	9.8	15.1	13	1	ABF22178	Oligonucleotide SE
419	9.8	15.1	13	1	ABF32116	Oligonucleotide SE
C 420	9.8	15.1	13	1	ABF39905	Oligonucleotide SE
C 421	9.8	15.1	13	1	ABF73693	Oligonucleotide SE
C 422	9.8	15.1	13	1	ABH33543	Oligonucleotide SE
423	9.8	15.1	13	1	ABH10962	Oligonucleotide SE
C 424	9.8	15.1	13	1	ABF22973	Oligonucleotide SE
C 425	9.8	15.1	13	1	ABH15571	Oligonucleotide SE
426	9.8	15.1	13	1	ABH44502	Oligonucleotide SE
427	9.8	15.1	13	1	ABH62561	Oligonucleotide SE
428	9.8	15.1	13	1	ABC73212	Oligonucleotide SE
429	9.8	15.1	13	1	ABC01962	Oligonucleotide SE
C 430	9.8	15.1	13	1	ABC60969	Oligonucleotide SE
431	9.8	15.1	13	1	ABC61676	Oligonucleotide SE
C 432	9.8	15.1	13	1	ABF31799	Oligonucleotide SE
433	9.8	15.1	13	1	ABF32795	Oligonucleotide SE
C 434	9.8	15.1	13	1	ABH22410	Oligonucleotide SE
C 435	9.8	15.1	13	1	ABH05561	Oligonucleotide SE
436	9.8	15.1	13	1	ABH34279	Oligonucleotide SE
C 437	9.8	15.1	13	1	ABH16293	Oligonucleotide SE
C 438	9.8	15.1	13	1	ABH44503	Oligonucleotide SE
C 439	9.8	15.1	13	1	ABC73213	Oligonucleotide SE
C 440	9.8	15.1	13	1	ABC99215	Oligonucleotide SE
441	9.8	15.1	13	1	ABF04646	Oligonucleotide SE
442	9.8	15.1	13	1	ABC55719	Oligonucleotide SE
C 443	9.8	15.1	13	1	ABF09481	Oligonucleotide SE
444	9.8	15.1	13	1	ABC16848	Oligonucleotide SE
445	9.8	15.1	13	1	ABF17417	Oligonucleotide SE
C 446	9.8	15.1	13	1	ABF32117	Oligonucleotide SE
C 447	9.8	15.1	13	1	ABH21397	Oligonucleotide SE
C 448	9.8	15.1	13	1	ABF75985	Oligonucleotide SE
C 449	9.8	15.1	13	1	ABH38625	Oligonucleotide SE
C 450	9.8	15.1	13	1	ABC68819	Oligonucleotide SE
451	9.8	15.1	13	1	ABC44906	Oligonucleotide SE
452	9.8	15.1	13	1	ABC74670	Oligonucleotide SE
453	9.8	15.1	13	1	ABF04648	Oligonucleotide SE
454	9.8	15.1	13	1	ABC05436	Oligonucleotide SE
C 455	9.8	15.1	13	1	ABC05437	Oligonucleotide SE
C 456	9.8	15.1	13	1	ABC06575	Oligonucleotide SE
C 457	9.8	15.1	13	1	ABC58519	Oligonucleotide SE
458	9.8	15.1	13	1	ABF09480	Oligonucleotide SE
C 459	9.8	15.1	13	1	ABF76689	Oligonucleotide SE
460	9.8	15.1	13	1	ABF67882	Oligonucleotide SE
C 461	9.8	15.1	13	1	ABF98633	Oligonucleotide SE
462	9.8	15.1	13	1	ABF73728	Oligonucleotide SE
463	9.8	15.1	13	1	ABC92814	Oligonucleotide SE
C 464	9.8	15.1	13	1	ABC20850	Oligonucleotide SE
465	9.8	15.1	13	1	ABC73225	Oligonucleotide SE
466	9.8	15.1	13	1	ABF31798	Oligonucleotide SE
C 467	9.8	15.1	13	1	ABF32794	Oligonucleotide SE
C 468	9.8	15.1	13	1	ABF33181	Oligonucleotide SE
469	9.8	15.1	13	1	ABF75984	Oligonucleotide SE
C 470	9.8	15.1	13	1	ABF58450	Oligonucleotide SE
471	9.8	15.1	13	1	ABF90374	Oligonucleotide SE

C 472	9.8	15.1	13	1	ABF66699	Oligonucleotide SE
473	9.8	15.1	13	1	ABH57530	Oligonucleotide SE
C 474	9.8	15.1	13	1	ABC75154	Oligonucleotide SE
C 475	9.8	15.1	13	1	ABF05601	Oligonucleotide SE
C 476	9.8	15.1	13	1	ABC55718	Oligonucleotide SE
C 477	9.8	15.1	13	1	ABF17416	Oligonucleotide SE
478	9.8	15.1	13	1	ABF31794	Oligonucleotide SE
C 479	9.8	15.1	13	1	ABF31795	Oligonucleotide SE
480	9.8	15.1	13	1	ABF3904	Oligonucleotide SE
C 481	9.8	15.1	13	1	ABH00239	Oligonucleotide SE
482	9.8	15.1	13	1	ABF50558	Oligonucleotide SE
C 483	9.8	15.1	13	1	ABH12784	Oligonucleotide SE
C 484	9.8	15.1	13	1	ABF88569	Oligonucleotide SE
C 485	9.8	15.1	13	1	ABF65173	Oligonucleotide SE
486	9.8	15.1	13	1	ABH15570	Oligonucleotide SE
487	9.8	15.1	13	1	ABH16218	Oligonucleotide SE
488	9.8	15.1	13	1	ABH16292	Oligonucleotide SE
C 489	9.8	15.1	13	1	ABH42927	Oligonucleotide SE
C 490	9.8	15.1	13	1	ABH62560	Oligonucleotide SE
491	9.8	15.1	13	1	ABC68818	Oligonucleotide SE
492	9.8	15.1	13	1	ABC34456	Oligonucleotide SE
493	9.8	15.1	13	1	ABH21396	Oligonucleotide SE
C 494	9.8	15.1	13	1	ABH22411	Oligonucleotide SE
495	9.8	15.1	13	1	ABF75610	Oligonucleotide SE
C 496	9.8	15.1	13	1	ABF75611	Oligonucleotide SE
497	9.8	15.1	13	1	ABH04580	Oligonucleotide SE
498	9.8	15.1	13	1	ABH12785	Oligonucleotide SE
C 499	9.8	15.1	13	1	ABC68985	Oligonucleotide SE
500	9.8	15.1	13	1	ABC73224	Oligonucleotide SE
501	9.8	15.1	13	1	ABC54294	Oligonucleotide SE
C 502	9.8	15.1	13	1	ABF67883	Oligonucleotide SE
C 503	9.8	15.1	13	1	ABF73729	Oligonucleotide SE
C 504	9.8	15.1	13	1	ABH00235	Oligonucleotide SE
C 505	9.8	15.1	13	1	ABF50559	Oligonucleotide SE
506	9.8	15.1	13	1	ABH29612	Oligonucleotide SE
C 507	9.8	15.1	13	1	ABF96467	Oligonucleotide SE
508	9.8	15.1	13	1	ABF62972	Oligonucleotide SE
509	9.8	15.1	13	1	ABC68984	Oligonucleotide SE
510	9.8	15.1	13	1	ABC99662	Oligonucleotide SE
511	9.8	15.1	13	1	ABC06245	Oligonucleotide SE
512	9.8	15.1	13	1	ABF07728	Oligonucleotide SE
C 513	9.8	15.1	13	1	ABC61677	Oligonucleotide SE
C 514	9.8	15.1	13	1	ABF35263	Oligonucleotide SE
515	9.8	15.1	13	1	ABF47356	Oligonucleotide SE
516	9.8	15.1	13	1	ABH33026	Oligonucleotide SE
517	9.8	15.1	13	1	ABF88568	Oligonucleotide SE
C 518	9.8	15.1	13	1	ABC20851	Oligonucleotide SE
519	9.8	15.1	13	1	ABC99214	Oligonucleotide SE
C 520	9.8	15.1	13	1	ABC99663	Oligonucleotide SE
C 521	9.8	15.1	13	1	ABC01963	Oligonucleotide SE
522	9.8	15.1	13	1	ABC37878	Oligonucleotide SE
C 523	9.8	15.1	13	1	ABC63814	Oligonucleotide SE
C 524	9.8	15.1	13	1	ABF66687	Oligonucleotide SE
C 525	9.8	15.1	13	1	ABF22179	Oligonucleotide SE
C 526	9.8	15.1	13	1	ABF72760	Oligonucleotide SE
C 527	9.8	15.1	13	1	ABF72761	Oligonucleotide SE
C 528	9.8	15.1	13	1	ABF98632	Oligonucleotide SE
C 529	9.8	15.1	13	1	ABF90375	Oligonucleotide SE
C 530	9.8	15.1	13	1	ABH15567	Oligonucleotide SE
531	9.8	15.1	13	1	ABH1661	Oligonucleotide SE
532	9.8	15.1	13	1	ABF66699	Oligonucleotide SE
C 533	9.8	15.1	13	1	ABH57531	Oligonucleotide SE
534	9.8	15.1	13	1	ABH57828	Oligonucleotide SE

ALIGNMENTS

RESULT 1  
 ADC02407/c  
 ID ADC02407 standard; DNA; 24 BP.  
 XX  
 AC ADC02407;